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(54) Title: GENE THERAPY METHOD USING FGF-5

(57) Abstract

A method is described for introducing an FGF-5 nucleic acid sequence into a mammalian host cell. The FGF-5 nucleic acid sequence lacks the signal sequence so that cells that are transformed with the sequence will not become tumirogenic. It is intended that the FGF-5 sequence is introduced into mammalian cells to promote angiogenesis. Preferably, the FGF-5 sequence is introduced into a human patient to treat myocardial ischemia or peripheral vascular disease.

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GENE THERAPY METHOD USING FGF-5

Field of the Invention

The present invention is in the field of gene therapy. More specifically, the present invention is in the field of gene therapy using the FGF-5 gene.

Background of the Invention

Fibroblast growth factors (FGFs) comprise a family of proteins with related amino acid structure. They are encoded by distinct genes and share sequence homology. Even though there are more than five FGFs, FGFs 1-5 will be discussed here. For example, FGF-1 is acidic FGF, FGF-2 is basic FGF, FGF-3 is int-2, FGF-4 is KFGF or HST, and FGF-5 is described herein.

The FGF-5 of the present invention was originally isolated as an oncogene. See Goldfarb et al. U.S. Patent Nos. 5,155,217 and 5,238,916, Zhan et al. "Human Oncogene Detected by a Defined Medium Culture Assay" (Oncogene (1987) 1:369-376), Zhan et al. "The Human FGF-5 Oncogene Encodes a Novel Protein Related to Fibroblastic Growth Factors" (Molecular and Cellular Biology (1988) 8:3487-3495), and Bates et al. "Biosynthesis of Human Fibroblast Growth Factor 5" (Molecular and Cellular Biology, (1991) 11:1840-1845). The disclosure of each of these patents and articles is hereby incorporated by reference in their entireties. The FGF-5 oncogene nucleic acid sequence was reported in both Goldfarb patents and in the Zahn et al. article (1988, 8). As discussed in each of these references, the FGF-5 gene is an oncogene which can transform cells into a tumorigenic state. Additionally, reports in the literature show that the related genes int-2 and HST can transform cells to be tumorigenic. See Theillet et al., Oncogene (1989) 4:915-922, and Goldfarb et al., Oncogene (1991 6:65-71.

Consequently, it is the aim of the present invention to use the FGF gene in gene therapy with human patients, while removing the oncogenic potential of this gene.

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Summary of the Invention

The present invention relates to a method for expressing FGF-5 in vivo, comprising introducing a nucleic acid sequence encoding FGF-5, without a signal sequence, into a vector that can infect mammalian cells and cause these cells to express FGF-5 without causing the cells to become tumorigenic.

The present inventor has discovered that to use FGF-5 in a gene therapy model in human patients, one must remove the signal sequence before administering the gene. Otherwise, the gene therapy may transform normal human cells into tumorigenic cells, which is obviously undesirable.

More specifically, the present invention relates to a gene therapy method for introducing an FGF-5 gene into a human cell of a patient suffering from myocardial ischemia or peripheral vascular disease comprising:

constructing a retroviral vector having a nucleic acid sequence encoding FGF-5, without a signal sequence, having an N terminus of GGGAGAAGCG TCTCGCCCCC AAAG (SEQ ID NO: 1), in operable linkage with the appropriate regulatory elements necessary to express the FGF-5 nucleic acid sequence in a human cell, to form the FGF-5 protein; and

introducing the vector into a cellular area in the human patient which is in need of treatment with the FGF-5 protein.

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Brief Description of the Drawings

Figure 1-A and Figure 1-B are the nucleic acid sequence for the FGF-5 gene which includes the signal sequence.

Figure 2 is the amino acid sequence for the FGF-5 gene which includes the signal sequence.

Figure 3 is the nucleic acid sequence for the FGF-5 gene starting at the 22nd amino acid of the sequence of Figure 1.

Figure 4 is the amino acid sequence for the FGF-5 gene starting at the 22nd amino acid of the sequence of Figure 2.

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Detailed Description of the Invention

As shown in Goldfarb et al. (U.S. Patent No. 5,155,217, the disclosure of which is hereby incorporated by reference in its entirety), FGFs 1-5 share a sequence homology between 41 and 50%. For example, column 9 of Goldfarb (*217) shows that there is 45% sequence identity between FGF-5 and basic FGF, 41% sequence homology between FGF-5 and acidic FGF, 52% sequence homology between FGF-5 and KFGF (also called HST), and 50% sequence homology between FGF-5 and int-2 (Goldfarb has used the designation FGF-3 throughout '217 but later changed the identity of their protein to FGF-5). See also Goldfarb et al. U.S. Patent No. 5,238,916. Basic FGF is more fully in U.S. Patent No. 5,155,214; 4,994,559; 5,401,701; and 5,439,818. Acidic FGF is disclosed in U.S. Patent No. 5,312,911. The disclosures of all of the U.S. patents listed above are hereby incorporated by reference in their entireties.

The FGF-5 protein has been shown to be synthesized *in vitro* in animal cells to yield a 29,500-dalton protein which was a secreted from tumor cells as a glycoprotein containing heterogeneous amounts of sialic acid. Glycosidase treatment suggested that FGF-5 has both N-linked and O-linked sugars. See Bates *et al.* "Biosynthesis of Human Fibroblast Growth Factor 5" (*Molecular and Cellular Biology*, (1991) 11:1840-1845), hereby incorporated by reference in its entirety.

The present invention describes the use of the FGF-5 nucleic acid sequence in a gene therapy method whereby the FGF-5 sequence is converted from an oncogene to a protooncogene (non tumorigenic) before it is introduced into human cells. As described above, the gene sequences are disclosed in the two Goldfarb patents ('217 and '916) and Zahn et al. "The Human FGF-5 Oncogene Encodes a Novel Protein Related to Fibroblastic Growth Factors" (Molecular and Cellular Biology (1988) 8:3487-3495), which are all hereby incorporated by reference in their entireties.

The FGF-5 oncogene is a 267 amino acid protein as compared to int-2, which is 240, HSTKS3, which is 206, and acidic and basic FGFs which are both 155 amino acids long. See Figures 1 and 2 for the nucleic acid and amino acid sequences of FGF-5, including the signal sequence. As stated above, the signal sequence of the FGF-5 oncogene must be removed before incorporating it into a gene therapy vector for human

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use. It is acceptible if enough of the signal sequence is removed so that the tumorigenic properties are eliminated from the FGF-5 molecule described in the Golfarb patent. Preferably, between 10 and 30 amino acids are removed from the N-terminus. More preferably, between 15 and 25 amino acids are removed from the N-terminus. Most preferably, the first 22 amino acids are removed from the N-terminus. (See Clements et al., Oncogene (1993) 8:1311-1316 which is hereby incorporated by reference in its entirety). See Figures 3 and 4 for the FGF-5 nucleic acid and amino acid sequences which begin at the 22nd amino acid of the sequences shown in Figures 1 and 2. Clements et al. disclose prokaryotic expression of the mature form of FGF-5 and describe silent mutations in the 5' end of the cDNA insert that increase the expression levels of FGF-5. The FGF-5 molecule of the present invention preferably contains those mutations.

The present gene therapy method of delivering FGF-5 to local areas in human patients is useful to treat human diseases of the vascular system, as well as enhancing the ability of neural cells to proliferate and for bone growth. See Morrison et al. "Basic 15 Fibroblast Growth Factor supports the survival of cerebral cortical neurons and primary culture" Proc. Natl. Acad. Sci. (USA) (1986) 83:7537-7541, which are hereby incorporated by reference in their entireties. There is also evidence that FGF-5 is a major musclederived survival factor for cultured spinal motor neurons (Hughes et al., Neuron (1993) 10:369-367), that FGF-5 is present in adult mouse central nervous system (Haub et al. 20 Proc. Natl. Acad. Sci. (USA) (1990) 87:8022-8026), that FGF-5 is a regulator of the hair growth cycle (Hebert et al., Cell (1984) 78:1017-1025), that FGF-5 promotes differentiation of cultured rat neurons (Lindholm et al., European Journal of Neuroscience (1994) 6:244-252), that FGF-5 may play a role in limbic system function or dysfunction (Gomez-Pinilla et al., Brain Research (1993) 606:79-86), that FGF-5 can play a role in the 25 biology of the outer retina (Bost et al., Exp. I. Res. (1992) 55:727-34), that basic FGF can ameliorate learning deficits in basal forebrain-lesioned mice (Ishihara et al., Jpn. J. Pharmacol. (1992) 59:7-13). Fibroblasts that have been engineered to express bFGF without a signal sequences have a more robust effect on the viability and function of grafted dopaminergic neurons than with fibroblasts that express bFGF with a signal 30 sequence (see Takayama et al., Nature Medicine (1995) 1:53-58). bFGF appears to be

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neuroprotective and neurotrophic (see Cheng and Mattson, Neuron (1991) 7:1031-1041; Freese et al., Brain Research (1992) 575:351-355; Finkelstein et al., Stroke (1993) 24 (supp. 1):141-143) angiogenic (Baffour et al., Jour. Vasc. Surg. (1992) 16:181-191); and osteogenic (Kawaguchi et al., Endocrinology (1993) 135:774-781; Nagai et al., Bone (1995) 16:367-373; Nakamura et al., Endocrinology (1995) 136:1276-1284; and Mayahara et al., Growth Factors (1993) 9:73-80). Also, it is contemplated that the FGF-5 gene will be useful for many of the uses shown for other FGFs. Accordingly, delivery of the FGF-5 gene will be useful in a variety of vascular, cardiovascular, neuronal, osteogenic, and other indications to correct or regulate cellular dysfunction. Preferably, the FGF-5 gene administered for angiogenic uses or to support their growth and/or proliferation or neuronal cells. More preferably, the FGF-5 nucleic acid sequence is administered to promote blood vessel growth in myocardial ischemia.

Definitions

The term "polynucleotide" or "nucleic acid sequence" as used herein refers to a polymer of nucleotides of any length, preferably deoxyribonucleotides, and is used interchangeably herein with the terms "oligonucleotide" and "oligomer." The term refers only to the primary structure of the molecule. Thus, this term includes double- and single-stranded DNA, as well as antisense polynucleotides. It also includes known types of modifications, for example, the presence of labels which are known in the art, methylation, end "caps," substitution of one or more of the naturally occurring nucleotides with an analog, internucleotide modifications such as, for example, replacement with certain types of uncharged linkages (e.g., methyl phosphonates, phosphotriesters, phosphoamidates, carbamates, etc.) or charged linkages (e.g., phosphorothioates, phosphorodithioates, etc.), introduction of pendant moieties, such as, for example, proteins (including nucleases, toxins, antibodies, signal peptides, poly-L-lysine, etc.), intercalators (e.g., acridine, psoralen, etc.), chelators (e.g., metals, radioactive species, boron, oxidative moieties, etc.), alkylators (e.g., alpha anomeric nucleic acids, etc.). The term "gene" is used to describe the coding sequence for the polypeptide of interest, for example, FGF-5.

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By "genomic" is meant a collection or library of DNA molecules which correspond to the sequence found in chromosomal DNA as opposed to spliced mRNA. By "cDNA" is meant a DNA sequence that hybridizes to a complimentary strand of mRNA.

"Regulatory" or "control sequence" refers to polynucleotide sequences which are necessary to effect the expression of coding sequences to which they are ligated. The nature of such control sequences differs depending upon the host organism; in eukaryotes, generally, such control sequences include promoters and transcription termination sequences. The term "control sequences" is intended to include, at a minimum, all components whose presence is necessary for expression, and may also include additional components whose presence is advantageous, for example, leader sequences and fusion partner sequences.

"Operably linked" refers to a juxtaposition wherein the components so described are in a relationship permitting them to function in their intended manner. A control sequence "operably linked" to a coding sequence so that expression of the coding sequence is achieved under conditions compatible with the control sequences.

A "vector" or "plasmid" is a nucleic acid sequence in which another polynucleotide segment is attached, so as to bring about the replication and/or expression of the attached segment in a host cell. Vectors are used routinly in recombinant DNA techniques. Any extrachromosomal small genome such as a plasmid, phage, or virus is a potential vector.

"Retroviral vector" is a vector derived from a retrovirus and it has the capability to insert a gene or DNA fragment into the host chromosomal genome by a recombinational event, so that the DNA fragment can be expressed in a host cell. See Singer, M. and Berg, P., Genes and Genomes, Mill Valley, CA (1991) pp. 310-314, which is hereby incorporated by reference. Retroviruses are RNA viruses (the viral genome is RNA). The genomic RNA is reverse transcribed into DNA after it enters the cell and then it is integrated stably and efficiently into the chromosomal DNA of transduced cells. See Mulligan, R.C., In: Experimental Manipulation of Gene

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Expression, M. Inouye (ed), 155-173 (1983); Mann, R. et al., Cell (1983) 33:153-159; Cone, R.D. and R. C. Mulligan, Proc. Natl. Acad. Sci. (USA), (1984) 81:6349-6353 which are hereby incorporated by reference in their entireties.

"Transformation", as used herein, refers to the insertion of an exogenous polynucleotide into a host cell, irrespective of the method used for the insertion, for example, direct uptake, particle mediated, transduction, f-mating or electroporation. The exogenous polynucleotide may be maintained as a non-integrated vector, for example, a plasmid, or alternatively, may be integrated into the host genome. Examples of particle mediated transduction are shown in U.S. Patent Nos. 4,945,050 and 5,149,655, which are hereby incorporated by reference in their entireties.

"Homology" refers to the degree of similarity between x and y. The correspondence between the sequence from one form to another can be determined by techniques known in the art. For example, they can be determined by a direct comparison of the sequence information of the polynucleotide. Alternatively, homology can be determined by hybridization of the polynucleotides under conditions which form stable duplexes between homologous regions (for example, those which would be used prior to S₁ digestion), followed by digestion with single-stranded specific nuclease(s), followed by size determination of the digested fragments.

As used herein, x is "heterologous" with respect to y if x is not naturally associated with y in the identical manner; i.e., x is not associated with y in nature or x is not associated with y in the same manner as is found in nature.

As used herein, the term "protein" or "polypeptide" refers to a polymer of amino acids and does not refer to a specific length of the product; thus, peptides, oligopeptides, polypeptides, proteins, and polyproteins, as well as fragments of these, are included within this definition. This term also does not refer to, or exclude, post expression modifications of the protein, for example, glycosylations, acetylations, phosphorylations and the like. Included within the definition are, for example, proteins containing one or more analogs of an amino acid (including, for example, unnatural amino acids, etc.),

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proteins with substituted linkages, as well as other modifications known in the art, both naturally occurring and non-naturally occurring.

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A polypeptide or protein or amino acid sequence "derived from" or "coded by" a designated nucleic acid sequence refers to a polypeptide having an amino acid sequence identical to that of a polypeptide encoded in the sequence, or a portion thereof wherein the portion consists of at least 3-5 amino acids, and more preferably at least 8-10 amino acids, and even more preferably at least 11-15 amino acids, or which is immunologically identifiable with a polypeptide encoded in the sequence. This terminology also includes a polypeptide expressed from a designated nucleic acid sequence.

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"Alleles" and "variants" refers to a polypeptide that differs from the native specified protein by virtue of one or more amino acid substitutions, deletions, or insertions. The amino acid substitutions can be conservative amino acid substitutions or substitutions to eliminate non-essential amino acid residues such as to alter a glycosylation site, a phosphorylation site, an acetylation site, or to alter the folding pattern by altering the position of the cysteine residue that is not necessary for function, etc. Conservative amino acid substitutions are those that preserve the general charge, hydrophobicity/hydrophilicity and/or steric bulk of the amino acid substituted, for example, substitutions between the members of the following groups are conservative substitutions: Gly/Ala, Val/Ile/Leu, Asp/Glu, Lys/Arg, Asn/Gln, and Phe/Trp/Tyr.

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"Signal sequence" is used to describe the N-Terminal amino acids that enable the polypeptide to be transported outside the boundaries of the cells in which it is made. As stated above, it is this sequence that enables the FGF-5 nucleic acid sequence to transform cells into a tumorigenic state. In FGF-5, it is the first 59 or, more preferably, the first 61 amino acids at the N-Terminus that constitute the signal sequence.

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The term "cardiovascular indication" as used herein refers to a diagnosis or presumptive diagnosis of cardiovascular disease or conditions affecting the heart that are associated with atheroscerosis, ischemic syndromes, cardiomyopathies, arrhythmias, dysrrhythmias, hypertension and infections. The diagnosis can be made based on pain, fatigability, weakness, palpitations, and systemic symptoms that may be due to the cardiac disease or that may accompany it. Determination of a cardiovascular indication

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may include a physical exam and other non-invasive diagnostic procedures including radionuclide imaging, positon emission tomography, magnetic resonance imaging, echocardiography, and can also include venous and arterial cannulation and pulmonary and cardiac catheterization used in diagnosis of the cardiac condition.

The term "administering to intrapericardially" or "administering into the pericardial space" as used herein refers to any method of administration that effects delivery of a therapeutic agent into the pericardial space. The pericardial space may be the entire region comprising the pericardial space, or only a part of it. The term "administering into pericardial space" is synonymous with the terms "intrapericardial delivery" and "pericardial delivery", and can include delivery to subregions of the pericardial space that form interfaces between the pericardial space and the tissue that surrounds and forms it. The administration into pericardial space can be accomplished by, for example, the following means of administration including injection, laser, catheter, pump. Intrapericardial delivery of the polynucleotides and the drugs of the invention can be accomplished by the methods of such delivery as disclosed in, for example, U.S. Patent Nos. 5,137,510, 5,269,326, and 5,213,570, herein incorporated by reference.

Vectors and Expression Systems

The following expression systems describe vectors, promoters and regulatory elements that are useful for gene therapy applications for the delivery of the FGF-5 polynucleotide. Vectors and expression systems useful for the present invention include viral and non-viral systems. Example viral delivery systems include retroviruses, adenoviruses, adeno-associated viruses (AAV), sindbis and herpes viruses. In one aspect of the present invention, the viral vector is capable of integrating the FGF-5 nucleic acid sequence into the host cell genome for long term expression. Examples of vectors that can integrate in this fashion are retroviruses and AAV. One preferred retrovirus is a murine leukemia virus. However, it may be preferred to avoid integration into the host cell genome. For example, when short term administration of FGF-5 is

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required, long term expression can be unecessary and possibly undesireable.

Non-viral vectors include naked DNA and DNA formulated with cationic lipids or liposomes.

Preferably, the FGF-5 nucleic acid coding sequence is administered in one of the above systems to a patient's cells without the signal sequence. The description below is directed to means for including the FGF-5 coding sequence in a larger sequence that will facilitate expression of the FGF-5 polypeptide.

Retroviral vectors are produced by genetically manipulating retroviruses. Retroviral vectors are effective for integration into the host cell genome, as explained above. However, they only infect dividing cells. Retroviral vectors contain RNA. In the present invention the viral RNA vector contains the FGF-5 gene, and once it enters the cell, it is reverse transcribed into DNA and stably integrated into the host cell genome.

The wild type retrovirus genome contains three genes: the gag, pol, and env genes, which are flanked by the long terminal repeat (LTR) sequences. The gag gene encodes the nucleocapsid proteins, the pol gene encodes the viral enzymes including reverse transcriptase and integrase, the env gene encodes the viral envelope glycoproteins. The 5' and 3' LTRs serve to promote transcription and polyadenylation of virion RNAs. Adjacent to the 5' LTR are sequences necessary for reverse transription of the genome (the tRNA primer binding site) and for efficient encapsulation of viral RNA into particles (the Psi site). See Mulligan, R.C., In: Experimental Manipulation of Gene Expression, M. Inouye (ed), 155-173 (1983); Mann, R. et al., Cell (1983) 33:153-159; Cone, R.D. and R. C. Mulligan, Proc. Natl. Acad. Sci. (USA), (1984) 81:6349-6353.

More specifically, the present invention contemplates constructing a vector in which the gag, pol, and env genes are removed and replaced with the FGF-5 gene. The LTR, psi sequence and primer binding sites are also present to facilitate vector replication. The vector is transformed into a packaging animal cell line which contains the gene sequences for the gag, pol, and env genes in its genome and which constitutively express those proteins. These proteins are

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usually expressed from a heterologous promoter (eg. CMV) and the genes are not operably linked to sequences (such as psi, LTR which are required for viral replication). This cell will make empty viral particles and is a recipient for the vector described above which contains the FGF-5 gene, the psi and primer binding sequences as well as the LTR sequences. The cell can be transiently transfected with the vector to produce the product (viral particle with the FGF-5 vector). Preferably, the product virions are used to infect a second packaging cell line which then can permanently produces the viral particles.

The retroviral vector can be packaged by transfecting the FGF-5 nucleic acid sequence into cells expressing the gag-pol and env genes. These "packaging cell lines" are mammalian tissue culture cell lines which express structural proteins of a retrovirus and produce retrovirus-like particles. They are ncapable of producing infectious virions. Transfecting retroviral vectors (with the FGF-5 nucleic acid sequence) into packaging cell lines results in the production of retroviral vector particles with the desired genetic construction. Packaging cell lines are publically available and include Crip, GPE86, PA317, and PG13. See Miller et al., J. Virol.(1991) 65:2220-2224, Cone et al., Proc. Natl. Acad. Sci. (USA), (1988) 85:6460-6464, Eglitis et al., Biotechniques (1988) 4:608-614, Miller et al., Human Gene Ther. (1990) 1:5-14, which are all hereby incorporated by reference in their entireties.

Also, AAV are advantageous because they replicate to a high titer, they integrate efficiently, are not pathogenic to humans, are stable, easy to purify, and they infect non-dividing cells. An AAV vector is constucted by inserting the FGF-5 coding sequence, under the control of a suitable promoter/enhancer, between the AAV LTRs, which are the only sequences required in cis for AAV replication. This DNA construct is transfected into a suitable human cell line in the presence of another plasmid which expresses Rep and CAP, the AAV coding regions needed for replication. At a suitable time post-transfection, the cells are infected with a helper virus, suhe as Adenovirus or Herpes Simplex virus. After infection, vector particles carrying the FGF-5 gene are harvested from these

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cells. The AAV particles are purified from contaminating Adenovirus or Herpes Virus by standard protocols.

Adeno virus is advantageous because it infects a wide variety of cells, infects non-dividing cells, produces a high titer, the biology is well understood, and it can accept large inserts. The adenovirus gene expression is controlled by a cascade of genes. For example, the gene expression order is "immediate early", "early", DNA synthesis, and late or structural genes. These genes are turned on in sequence. The master gene that is turned on first is E1A. One preferred embodiment would involve replacing the E1A gene with the FGF-5 gene and transfecting this vector into cells that constitutively produce E1A, such as 293 cells which are publically available. The vector contains all the genes necessary for virion production and the cell line provides the missing E1A protein. Consequently, the virion is produced which contains the FGF-5 sequence.

One non-viral system that can be used is the T7/T7 system. Here a short promoter sequence recognized by the bacterial virus T7 polymerase is placed on a vector upstream of the FGF-5 gene. The vector can then be inserted into cells and the missing T7 polymerase can be added to obtain gene transcription.

Alternatively, a vector containing the following sequences can be made, the T7 promoter sequence, the T7 polymerase gene, another copy of the T7 promoter sequence, and the FGF-5 gene. In this embodiment, the vector is transformed into cells and simply requires a small amount of T7 polymerase to initiate.

Therafter, the vector directs the manufacture of its own polymerase.

Although the methodology described is believed to contain sufficient details to enable one skilled in the art to practice the present invention, other items not specifically exemplified, such as plasmids, can be constructed and purified using standard recombinant DNA techniques described in, for example, Sambrook et al. (1989), MOLECULAR CLONING: A LABORATORY MANUAL, 2d edition (Cold Spring Harbor Press, Cold Spring Harbor, N.Y.), and Ausubel et al., CURRENT PROTOCOLS IN MOLECULAR BIOLOGY

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(1994), (Greene Publishing Associates and John Wiley & Sons, New York, N.Y.). under the current regulations described in United States Dept. of HEW, NATIONAL INSTITUTE OF HEALTH (NIH) GUIDELINES FOR RECOMBINANT DNA RESEARCH. These references include procedures for the following standard methods: cloning procedures with plasmids, transformation of host cells, cell culture, plasmid DNA purification, phenol extraction of DNA, ethanol precipitation of DNA, agarose gel electrophoresis, purification of DNA fragments from agarose gels, and restriction endonuclease and other DNA-modifying enzyme reactions.

Gene therapy can be practiced according to the invention by genes that are under regulatory control of appropriate regulatory sequences for transformation or infection of myocytes, cells within the pericardium, cells at the epicardium, or any cells in a region of the heart accessible to an intrapericardially delivered gene. When the genes are directed to nerve cells, the genes must be under the appropriate regulatory elements that enable expression in those cells. Gene therapy can be practiced as follows using coding regions for any therapeutic appropriate for treatment of a cardiovascular or neural indication.

As explained above, gene therapy strategies for delivery of the FGF-5 gene nucleic sequence can utilize viral or non-viral vector approaches in *in vivo* or *ex vivo* modality. Expression of such coding sequence can be induced using endogenous mammalian, viral or other heterologous promoters. Expression of the coding sequence *in vivo* can be either constitutive or regulated.

For delivery using viral vectors, any of a number of conventional viral vectors can be used, as described in Jolly, Cancer Gene Therapy (1994) 1:51-64. Promoters that are suitable for use with these vectors are also conventional in the art and include the Moloney retroviral LTR, CMV promoter and the mouse albumin promoter. Virus competent for one round of replication can be produced and injected directly into the animal or humans or by transduction of an autologous cell ex vivo, followed by injection in vivo as described in Zatloukal et al., Proc. Natl. Acad. Sci. USA (1994) 91:5148-5152.

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Delivery

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Preferably, the FGF-5 gene is administered to the local area of the pericardium or neural cells. More preferably, the FGF-5 gene is delivered to the pericardium without the signal sequence. The FGF-5 nucleic acid sequence may be delivered into the pericardial space by any method conventional in the art, such as that described in Barr et al., Gene Therapy (1994) 1:51-58. Barr et al. describe gene delivery via catheter-mediated infusion of replication defective adenovirus into the coronary arterial circulation. High level expression of an exogenous gene was obtained throughout the thickness of the ventricular and aterial walls within the distribution of the injected coronary artery. The FGF-5 nucleic acid sequence may be linked to tissue specific promoters or leader sequences for expression in cardiac muscle cells, for example, the untranslated leader sequence of dystrophin DNA, or regulatory regions in the muscle creatine kinase gene such as that described in Cox et al., Nature (1993) 364:725-729.

Delivery of genes to the intrapericardial space is a safer and more effective method of accomplishing myocardial gene therapy. Accordingly, delivery of genes to the pericardial space does not require mechanical violation of the myocardium as does direct myocardial injection. Because intrapericardially delivered agents have access to the entire myocardial surface the ease and effectiveness with which genes can be delivered to large areas of myocardium is increased. Access the coronary circulation causes perfusion of the entire heart with these agents. Also, the pericardium is more easily transducible than myocardium and, thus, that expression of gene products in the pericardial space retains access to myocardium.

Furthermore, the exposure time of nucleic acids and/or viruses to cells, which is an important determinant of transduction or infection efficiency, increases. Genetic agents deposited in the pericardial space are not subject to rapid dilution, drainage, or dissipation due to blood flow or lymphatic clearance, and thus have much longer exposure times than vascularly delivered agents, also increasing the transduction of infection efficiency of the genes. Such an advantage achieved by the method of the invention, translates into much higher transduction

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or infection efficiency with genes and/or viruses in either the myocardium or the pericardium than is achievable in the coronary vessel. Lastly, because pericardium is highly efficient at expressing certain proteins and in some cases is even more efficient than myocardium at this task, the method of the invention is a new and improved method of delivery of genes for gene therapy for treatment of a cardiovascular indication.

Practice of the invention also includes, for example, delivering the FGF-5 genes into the pericardial space, optionally in combination with cardiovascular therapeutics, in liposomal compositions, including heterovesicular liposomes. Delivery in liposomes increases the efficacy of the gene or cardiovascular therapeutics, reduces the dosage requirements and augments the benefits of any cardiovascular therapeutic delivered into the pericardial space.

Additionally, the FGF-5 gene can be delivered to nerve tissue. Actual delivery methods may vary, depending on the sites of the nerves to be affected. For example, administration to nerve tissue may be by encapsulating the FGF nucleotide sequence in a herpes virus which will specifically target nerve cells.

For in vivo therapy, the coding sequence can be delivered into the intrapericardial space by direct injection, or into pericaridial tissue by delivery such as, for example, those systems described in U.S. Patent Nos. 5,137,510, 5,213,570, and 5,269,326. Promoters suitable for use in this manner include endogenous and heterologous promoters such as those described herein. Any promoter appropriate for the expression of the gene selected for the therapy is contemplated by the method of the invention. The coding sequence can be injected in a formulation comprising a buffer that can stablize the coding sequence and facilitate transduction thereof into cells and/or provide targeting, as described in Zhu et al., Science (1993) 261:209-211.

Expression of the FGF-5 coding sequence in vivo (by either viral or non-viral vectors) can be regulated by use of regulated gene expression promoters as described in Gossen et al., Proc. Natl. Acad. Sci. (USA) (1992) 89:5547-5551. For example, the coding sequence selected for the therapy can be regulated by tetracycline responsive promoters. These promoters can be regulated in a positive or negative fashion by

treatment with the regulator molecule. Additionally, the FGF-5 gene may be introduced into cells under the control of promoters which are activated using radiotherapy. For example, U.S. Patent No. 5,205,152 entitled "Cloning and Expression of Early Growth Regulatory Protein Genes" shows that the Egr-1 gene is one of the best radiation induced genes and may be activated by exposure to radiation. WO 92/11033 disclosed genetic constructs which comprise an enhancer-promoter region which is responsive to radiation, and at least one structural gene whose expression is controlled by the enhancer-promoter. The U.S. Patent and the PCT application are hereby incorporated by reference in their entireties.

10 For non-viral delivery, the FGF-5 coding sequence can be inserted into conventional vectors that contain conventional control sequences for high level expression, and then incubated with synthetic gene transfer molecules such as polymeric DNA-binding cations like polylysine, protamine, and albumin, linked to cell targeting ligands such as asialoorosomucoid, as described in Wu and Wu, J. Biol. Chem. (1987) 15 262: 4429-4432; insulin, as described in Hucked et al., Biochem. Pharmacol. (1990) 40:253-263; galactose, as described in Plank et al., Bioconjugate Chem. (1992) 3:533-539; lactose, as described in Midoux et al., Nucleic Acids Res. (1993) 21:871-878; or transferrin, as described in Wagner et al., Proc. Natl. Acad. Sci. (USA) (1990) 87:3410-3414. Other delivery systems include the use of liposomes to encapsulate DNA 20 comprising the gene under the control of a variety of tissue-specific or ubiquitouslyactive promoters, as described in Nabel et al., Proc. Natl. Acad. Sci. (USA) (1993) 90:11307-11311, and Philip et al., Mol. Cell Biol. (1994) 14:2411-2418. Further nonviral delivery suitable for use includes mechanical delivery systems such as the biolistic approach, as described in Woffendin et al., Proc. Natl. Acad. Sci. (USA) (1994) 25 91(24):11581-11585. Moreover, the coding sequence and the product of expression of such can be delivered through deposition into the pericaridial space of photopolymerized hydrogel materials such as Focalgel[®]. Furthermore, the FGF-5 gene sequence can be inserted into a host cell by direct uptake or particle mediated transduction. The FGF-5 sequence may be maintained as a non-integrated vector, for example, a plasmid, or 30 alternatively, may be integrated into the host genome. Examples of particle mediated

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transduction are shown in U. S. Patent Nos. 4,945,050 and 5,149,655, which are hereby incorporated by reference in their entireties.

As stated above, naked DNA can be adminstered to muscle tissue. See Wolff, JA et al. entitled Direct gene transfer into mouse muscle in vivo, Science (1990) 247:1465-1468; Kitsis et al., Hormonal modulation of a gene injected into rat heart in vivo, Proc. Natl. Acad. Sci. (1991) 88:4138-4142; Lin et al., Expression of recombinant genes in myocardium in vivo after direct injection of DNA, Circulation (1990) 82:2217-2221; and Buttrick et al., Behavior of genes directly injected into rat heart in vivo, Circ. Res. (1992) 70:193-198. The above references are hereby incorporated by reference in their entireties.

To practice one aspect of the invention, the diagnosis of a cardiovascular condition is made, and the appropriate dosages are determined on the basis of the diagnosis. The invention is practiced to prevent, reduce or treat a cardiovascular condition.

The method of the invention applies to any cardiovascular indication, for example a diagnosis of: (1) atherosclerosis, and conditions that predispose one to pathological atherosclerotic plaque development in the coronary arteries including lipid/cholesterol deposition, macrophage/inflammatory cell recruitment, plaque rupture, thrombosis, platelet deposition, neointimal proliferation; (2) ischemic syndromes and attendent syndromes, including but not limited to myocardial infarction, stable and unstable angina, coronary artery restenosis following percutaneous transluminal, coronary angioplasty, reperfusion injury; (3) cardiomyopathies, including but not limited to cardiomyopathies caused by ischemic syndromes, cardiotoxins such as alcohol and chemotherapeutic agents like adriamycin, infections, such as viral, cytomegalovirus (CMV), and parasitic (trypanosoma cruzi), hypertension, metabolic diseases, (including but not limited to uremia, beriberi, glycogen storage disease), radiation, neuromuscular disease (such as Duchenne's muscular dystrophy), infiltrative diseases (including but not limited to sarcoidosis, hemochromatosis, amyloidosis, Fabry's disease, Hurler's syndrome), trauma, and idiopathic causes; (4) a/dysrrhythmias (including but not limited to a/dysrrhythmias resulting from the same causes listed above for cardiomyopathies);

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(5) infections (including bacterial, viral, fungal, and parasitic causes); (6) cardiac tumors; (7) inflammatory conditions (including but not limited to myocarditis, pericarditis, endocarditis, immune cardiac rejection and conditions resulting from idiopathic, autoimmune, or connective tissue diseases); and (8) hypertension.

The FGF-5 nucleotide sequence can be administered to the pericardial space and expressed in the heart tissue, including but not limited to, for example, pericardial tissue, myocardial tissue, epicardial tissue, or perivascular tissue. The sequence can be placed in a vector, such as a viral vector, or a plasmid vector. The polynucleotides may be presented into the pericardial space in any formulation commonly known in the art including buffers, excipients, gels, matrices and polymers. Appropriate formulations for the polynucleotides administered intrapericardially in the practice of the invention also include liposomal preparations such as, for example, those disclosed in U.S. Patent No. 5,422,120, WO 95/13796, WO 94/23697, WO 91/14445 and EP 524,968 B1, particularly including the heterovesicular liposomal preparations disclosed in these patents and applications.

The present invention will now be illustrated by reference to the following examples which set forth particularly advantageous embodiments. However, it should be noted that these embodiments are illustrative and are not to be construed as restricting the invention in any way.

20 Example 1

The coding sequence for FGF-5, without the signal sequence, is isolated by standard recombinant DNA techniques and placed in a retroviral vector and encapsulated in viral envelope for delivery intrapericardially. The retrovirus is delivered by laparoscopic cannulation or direct injection into the pericardial space of a patient who is suffering from myocardial ischemia or peripheral vascular disease. Alternatively, the coding sequence is placed in a plasmid vector and the vector is likewise delivered into the pericardial space. The coding sequences are linked with appropriate regulatory sequences and are delivered into the pericardial space by laparoscopic cannulation or direct injection. The FGF-5 nucleic acid sequence is expressed by the patient's cells in the

local area of the release and the FGF-5 protein induces the formation of new blood vessels.

The present invention has been described with reference to specific embodiments.

However, this application is intended to cover those changes and substitutions which may
be made by those skilled in the art without departing from the spirit and the scope of the appended claims.

SEQUENCE LISTING

GGGAGAAGCG	TCTCGCCCCC	AAAG	(SEQ	ID	NO:	1)
TTCTTCAGCC	ACCTGATCCT	CAGC	(SEQ	ID	NO:	2)
ATCCTCAGCG	CCTGGGCTCA	CGGG	(SEQ	ID	NO:	3)
CGTCTCGCCC	CCAAAGGGCA	ACCC	(SEQ	ID	NO:	4)

GGGCAACCCG GACCCGCTGC CACT (SEQ ID NO: 5)

CLAIM:

- 1. A method for expressing FGF-5 in vivo, comprising introducing a nucleic acid sequence encoding FGF-5, without a signal sequence, into a vector that can be introduced into mammalian cells to cause these cells to express FGF-5 without causing the cells to become tumorigenic.
- 2. A method in accordance with claim 1, wherein the FGF-5 sequence is administered to a human patient to induce angiogenesis in that patient.
- 3. A method in accordance with claim 1, wherein the vector is a retrovirus, an adenovirus, sindbis virus, herpes virus, or an adeno-associated virus.

 A method in accordance with claim 2 wherein the FGF-5 vector is used to treat myocardial ischemia or peripheral vascular disease.
- 4. A vector comprising a promoter operable in a eukaryotic cell, a nucleic acid sequence encoding FGF-5, without a signal sequence, the FGF-5 nucleic acid sequence being in operable linkage with the promoter.
- 5. A gene therapy method for introducing an FGF-5 gene into a human cell of a patient suffering from myocardial ischemia or peripheral vascular disease comprising:

constructing a retroviral vector having a nucleic acid sequence encoding FGF-5, without a signal sequence, having an N terminus of GGGAGAAGCG TCTCGCCCCC AAAG (SEQ ID NO: 1), in operable linkage with the appropriate regulatory elements necessary to express the FGF-5 nucleic acid sequence in a human cell, to form the FGF-5 protein; and

introducing the vector into a cellular area in the human patient which is in need of treatment with the FGF-5 protein.

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- 6. A method in accordance with claim 1, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of TTCTTCAGCC ACCTGATCCT CAGC (SEQ ID NO: 2).
- 7. A method in accordance with claim 1, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of ATCCTCAGCG CCTGGGCTCA CGGG (SEQ ID NO: 3).
- 8. A method in accordance with claim 1, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of GGGAGAAGCG TCTCGCCCCC AAAG (SEQ ID NO: 1).
- 9. A method in accordance with claim 1, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of CGTCTCGCCC CCAAAGGGCA ACCC (SEQ ID NO: 4).
- 10. A method in accordance with claim 1, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of GGGCAACCCG GACCCGCTGC CACT (SEQ ID NO: 5).
- 11. A method in accordance with claim 4, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of TTCTTCAGCC ACCTGATCCT CAGC (SEQ ID NO: 2).
- 12. A method in accordance with claim 4, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of ATCCTCAGCG CCTGGGCTCA CGGG (SEQ ID NO: 3).

- 13. A method in accordance with claim 4, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of GGGAGAAGCG TCTCGCCCCC AAAG (SEQ ID NO: SEQ ID NO: 1).
- 14. A method in accordance with claim 4, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of CGTCTCGCCC CCAAAGGGCA ACCC (ID SEQ ID NO: 4).
- 15. A method in accordance with claim 4, wherein the nucleic acid sequence encoding FGF-5 has an N terminus of GGGCAACCCG GACCCGCTGC CACT (SEQ ID NO: 5).

FIGURE 1

CCTCTCCCCT	TCTCTTCCCC	GAGGCTATGT	CCACCCGGTG	CGGCGAGGCG	GGCCAGAGCA
GAGGCACGCA	GCCGCACAGG	GGCTACAGAG	CCCAGAATCA	GCCCTACAAG	ATGCACTTAG
GACCCCCGCG	GCTGGAAGAA	TGAGCTTGTC	CTTCCTCCTC	CTCCTCTTCT	TCAGCCACCT
GATCCTCAGC	GCCTGGGCTC	ACGGGGAGAA	GCGTCTCGCC	CCCAAAGGGC	AACCCGGACC
CGCTGCCACT	GATAGGAACC	CTAGAGGCTC	CAGCAGCAGA	CAGAGCAGCA	GTAGCGCTAT
GTCTTCCTCT	TCTGCCTCCT	CCTCCCCCGC	AGCTTCTCTG	GGCAGCCAAG	GAAGTGGCTT
GGAGCAGAGC	AGTTTCCAGT	GGAGCCTCGG	GGCGCGGACC	GGCAGCCTCT	ACTGCAGAGT
GGGCATCGGT	TTCCATCTGC	AGATCTACCC	GGATGGCAAA	GTCAATGGAT	CCCACGAAGC
CAATATGTTA	AGTGTTTTGG	AAATATTTGC	TGTGTCTCAG	GGGATTGTAG	GAATACGAGG
AGTTTTCAGC	AACAAATTTT	TAGCGATGTC	AAAAAAAGGA	AAACTCCATG	CAAGTGCCAA
GTTCACAGAT	GACTGCAAGT	TCAGGGAGCG	TTTTCAAGAA	AATAGCTATA	ATACCTATGC
CTCAGCAATA	CATAGAACTG	AAAAAACAGG	GCGGGAGTGG	TATGTTGCCC	TGAATAAAAG
AGGAAAAGCC	AAACGAGGGT	GCAGCCCCCG	GGTTAAACCC	CAGCATATCT	CTACCCATTT
TCTTCCAAGA	TTCAAGCAGT	CGGAGCAGCC	AGAACTTTCT	TTCACGGTTA	CTGTTCCTGA
AAAGAAAAAT	CCACCTAGCC	CTATCAAGTC	AAAGATTCCC	CTTTCTGCAC	CTCGGAAAAA
TACCAACTCA	GTGAAATACA	GACTCAAGTT	TCGCTTTGGA	TAATATTAAT	CTTGGCCTTG
TGAGAAACCA	TTCTTTCCCC	TCAGGAGTTT	CTATAGGTGT	CTTCAGAGTT	CTGAAGAAAA
ATTACTGGAC	ACAGCTTCAG	CTATACTTAC	ACTGTATTGA	AGTCACGTCA	TTTGTTTCAG
TGTGACTGAA	ACAAAATGTT	TTTTGATAGG	AAGGAAACTG	(SEQ ID NO	: 6)

FIGURE 2

MSLSFLLLF FSHLILSAWA HGEKRLAPKG QPGPAATDRN PRGSSSRQSS SSAMSSSSAS SSPAASLGSQ GSGLEQSSFQ WSLGARTGSL YCRVGIGFHL QIYPDGKVNG SHEANMLSVL EIFAVSQGIV GIRGVFSNKF LAMSKKGKLH ASAKFTDDCK FRERFQENSY NTYASAIHRT EKTGREWYVA LNKRGKAKRG CSPRVKPQHI STHFLPRFKQ SEQPELSFTV TVPEKKNPPS PIKSKIPLSA PRKNTNSVKY RLKFRFG (SEQ ID NO: 7)

FIGURE 3

ATGAGCTTGT	CCTTCCTCCT	ССТССТСТТС	TTCAGCCACC	TGATCCTCAG	CGCCTGGGCT
CACGGGGAGA	AGCGTCTCGC	CCCCAAAGGG	CAACCCGGAC	CCGCTGCCAC	TGATAGGAAC
CCTAGAGGCT	CCAGCAGCAG	ACAGAGCAGC	AGTAGCGCTA	TGTCTTCCTC	TTCTGCCTCC
TCCTCCCCCG	CAGCTTCTCT	GGGCAGCCAA	GGAAGTGGCT	TGGAGCAGAG	CAGTTTCCAG
TGGAGCCTCG	GGGCGCGGAC	CGGCAGCCTC	TACTGCAGAG	TGGGCATCGG	TTTCCATCTG
CAGATCTACC	CGGATGGCAA	AGTCAATGGA	TCCCACGAAG	CCAATATGTT	AAGTGTTTTG
GAAATATTTG	CTGTGTCTCA	GGGGATTGTA	GGAATACGAG	GAGTTTTCAG	CAACAAATTT
TTAGCGATGT	CAAAAAAAGG	AAAACTCCAT	GCAAGTGCCA	AGTTCACAGA	TGACTGCAAG
TTCAGGGAGC	GTTTTCAAGA	AAATAGCTAT	AATACCTATG	CCTCAGCAAT	ACATAGAACT
GAAAAAACAG	GGCGGGAGTG	GTATGTTGCC	CTGAATAAAA	GAGGAAAAGC	CAAACGAGGG
TGCAGCCCCC	GGGTTAAACC	CCAGCATATC	TCTACCCATT	TTCTTCCAAG	ATTCAAGCAG
TCGGAGCAGC	CAGAACTTTC	TTTCACGGTT	ACTGTTCCTG	AAAAGAAAAA	TCCACCTAGC
CCTATCAAGT	CAAAGATTCC	CCTTTCTGCA	CCTCGGAAAA	ATACCAACTC	AGTGAAATAC
AGACTCAAGT	TTCGCTTTGG	A (SEQ ID	NO: 8)		

FIGURE 4

HGEKRLAPKG QPGPAATDRN PRGSSSRQSS SSAMSSSSAS SSPAASLGSQ GSGLEQSSFQ WSLGARTGSL YCRVGIGFHL QIYPDGKVNG SHEANMLSVL EIFAVSQGIV GIRGVFSNKF LAMSKKGKLH ASAKFTDDCK FRERFQENSY NTYASAIHRT EKTGREWYVA LNKRGKAKRG CSPRVKPQHI STHFLPRFKQ SEQPELSFTV TVPEKKNPPS PIKSKIPLSA PRKNTNSVKY RLKFRFG (SEQ ID NO: 9)

INTERNATIONAL SEARCH REPORT

International slication No PCT/US 97/02338

A. CLASSI IPC 6	FICATION OF SUBJECT MATTER C12N15/12 C07K14/50 C12N15/8	36 A61K48/00	
According to	o international Patent Classification (IPC) or to both national class	fication and IPC	
	SEARCHED		
IPC 6	ocumentation searched (classification system followed by classificat C12N C07K A61K		
Documentat	bon searched other than minimum documentation to the extent that	such documents are included in the fields s	earched
Electronic d	ists base consulted during the international search (name of data ba	se and, where practical, search terms used)	
C. DOCUM	MENTS CONSIDERED TO BE RELEVANT		
Category *	Citation of document, with indication, where appropriate, of the r	elevant passages	Relevant to claim No.
Υ .	JOURNAL OF INVESTIGATIVE MEDICINI vol. 43, no. SUPPL.2, 1995, page 278A XP002034595 GIORDANO F.J. ET AL.: "Reduced after recombinant Adenovirus med in-vivo fibroblast growth factor	ischemia iated	1-4
Α .	transfer." see abstract		5-15
Y	CANCER AND METASTASIS REVIEWS, vol. 9, 1990, pages 191-202, XP002034598 YAYON A. AND KLAGSBRUN M.: "Aut regulation of cell growth and transformation by basic fibrobla factor."		1-4
A	see abstract	-/	5-15
X Fur	ther documents are listed in the continuation of box C.	Patent family members are listed	in annex.
"A" docum "E" earlier filing "L" docum which citatic "O" docum other "P" docum later i	ment which may throw doubts on priority claim(s) or is cited to establish the publication date of another on or other special reason (as specified) ment referring to an oral disclosure, use, exhibition or means ment published prior to the international filing date but than the priority date claimed	T later document published after the into or priority date and not in conflict worked to understand the principle or to invention "X" document of particular relevance; the cannot be considered novel or cannot involve an inventive step when the desired to involve an inventive step when the cannot be considered to involve an in document is combined with one or in ments, such combination being obvious the art. "A" document member of the same patent.	th the application but hereby underlying the claimed invention to be considered to occurrent is taken alone claimed invention inventive step when the noise other such docupus to a person skilled it family
1	s actual completion of the international search 3 July 1997	Date of mailing of the international s	earu report
Name and	mailing address of the ISA European Patent Office, P.B. 5818 Patentiaan 2 NL - 2230 HV Ristorit Tel. (+ 31-70) 340-2040, Tx. 31 651 epo nl, Fax (+ 31-70) 340-3016	Authorized officer Mandl, B	

INTERNATIONAL SEARCH REPORT

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C.(Continue Category	citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.	
A	ONCOGENE, vol. 6, no. 8, 1991, pages 1437-1444, XP002034596 COULIER F. ET AL.: "Putative structure of the FGF6 gene product and role of signal peptide." see the whole document, especially from p.1441, last paragraph to p.1442 first paragraph and discussion	1-15	
A	MOLECULAR AND CELLULAR BIOLOGY, vol. 8, 1988, pages 3487-3495, XP002034597 ZHAN X. ET AL.: "The human FGF-5 oncogene encodes a novel protein related to fibroblast growth factors." cited in the application see the whole document	1-15	
A	CIRCULATION, vol. 84, no. 4, suppl., 1991, page II-420 (A1673) XP002034599 BARR E. ET AL.: "Induction of angiogenesis following in vivo gene transfer into myocardium." see abstract	1-15	

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International application No.

INTERNATIONAL SEARCH REPORT

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Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. X Claims Nos.: 1-3,3A,5-15 because they relate to subject matter not required to be searched by this Authority, namely: Remark: Although claims 1-3,3A,5-15 are directed to a method of treatment of (diagnostic method practised on) the human/animal body, the search has been carried out and based on the alleged effects of the compound/composition.
2. X Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically: After claim 3 and before calim 4, there is obviously another claim which has not got a number: "A method in accordance with claim 2" This claim was termed 3A. Claims 11-15, obviously refer to this claim 3A (not to claim 4]) and where searched as such.
3. Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
Remark on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.